Transformer-based generative model accelerating the development of novel BRAF inhibitors

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Table S1: Experimental results of the percentage of small molecules generated when GRUs of different layers are stacked.

	No. of layers =1	No. of layers =2	No. of layers =4
Invalid	4.368%±0.361%	4.104%±0.264%	4.270%±0.245%
Redundant	$0.038\% \pm 0.015\%$	0.034%±0.009%	0.038%±0.024%
Retained	95.594%±0.351%	95.862%±0.268%	95.692%±0.241%

Table S2: Test results of the predictive model on the PDBbind dataset.

	Number of samples	RMSE	R
Train	10680	1.034	0.85
Test	1186	1.463	0.68

Table S3: The percentage changes of molecules generated during reinforcement learning.

	Aver reward=10.0		Aver rew	ard=14.0	Aver reward=15.0		
	Transformer	GRU	Transformer	GRU	Transformer	GRU	
Invalid	1.080%±0.141%	3.508%±0.160%	1.483%±0.145%	2.772%±0.104%	1.170%±0.052%	2.690%±0.167%	
Redundant	14.473%±0.202%	9.584%±0.262%	19.247%±0.720%	20.626%±0.430%	30.273%±0.525%	33.828%±0.616%	
Retained	84.447%±0.097%	86.877%±0.502%	79.270%±0.740%	76.427%±0.429%	68.557%±0.552%	63.482%±0.456%	

Table S4: Details of the architecture and hyperparameters used for training the generative and predictive models.

	Generati	ve model	Transfer learning Reinforcement learning		ent learning	Predictive model	
Hyperparameter	Transformer- based model	GRU-based model	Transformer- based model	GRU-based model	Transformer- based model	GRU-based model	
Type of neural network	Transformer	RNN	Transformer	RNN	Transformer	RNN	RNN
No. of layers	12	1	12	1	12	1	3
Hidden size	768	1500	768	1500	768	1500	256
Stacked memory		1500		1500		1500	
Activation function	GELU	TanH	GELU	TanH	GELU	TanH	ReLU
Optimizer	AdamW	Adam	AdamW	Adam	AdamW	Adam	Adam

Loss function	Cross	Cross	Cross	Cross	Cross	Cross	MSELoss	
Loss function	Entropy Loss	WISELOSS						
Initial learning rate	0.00015	0.0005	0.00015	0.0005	0.00015	0.0005	0.001	
Batch size	48	128	48	128	50	50	48	
Gradient clipping	Yes							
No. of epochs	500	500	100	100	60	150	150	
GPU details	1 Tesla A100							
	GPU							
Implementation framework	PyTorch							

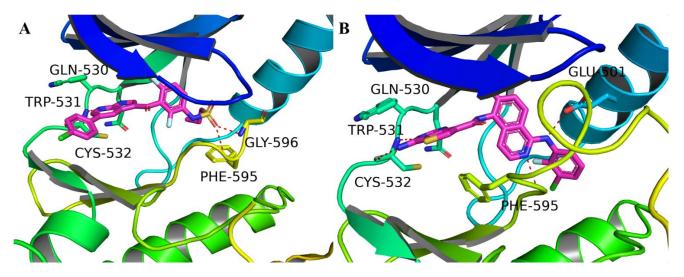


Figure S1. Two binding pockets of BRAF. A. the ATP binding pocket; B. the allosteric binding pocket.